

Amendments to the Specification:

Please replace the paragraphs at page 11, lines 15-19 with the following:

The EST and contig sequences regarding BIIA1 and BIIA2 are available through the Sanger Institute's sequencing genomics projects' Internet web page:-
www.sanger.ac.uk/projects/b-bovis/. The contig sequences regarding TIIA1 and TIIA2 are available through the NCBI BLAST server by selecting Apicomplexa from the United States' National Institutes of Health's National Library of Medicine's National Center for Biotechnology Information Eukaryotic genomes' BLAST Internet page:
http://www.ncbi.nlm.nih.gov/sutils/genom_tree.cgi?organism=euk.

In the Abstract of the Disclosure:

Please replace the Abstract with the attached Abstract provided herewith on a separate sheet.